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SEQUENCE LISTING

(1) GENERAL INFORMATION:

(i) APPLICANT: NG, Frank M.

JIANG, Woei-Jia

(ii) TITLE OF INVENTION: Treatment of Obesity

(iii) NUMBER OF SEQUENCES: 33

(iv) CORRESPONDENCE ADDRESS:

(A) ADDRESSEE: Metabolic Pharmaceuticals Ltd.

(B) STREET: 10 Wallace Avenue

(C) CITY: Toorak

(D) STATE: Victoria

(E) COUNTRY: Australia

(F) ZIP: 3142

(v) COMPUTER READABLE FORM:

(A) MEDIUM TYPE: Floppy disk

(B) COMPUTER: IBM PC compatible

(C) OPERATING SYSTEM: PC-DOS/MS-DOS

(D) SOFTWARE: PatentIn Release #1.0, Version #1.25

(vi) CURRENT APPLICATION DATA:

(A) APPLICATION NUMBER:

(B) FILING DATE:

(vii) PRIOR APPLICATION DATA:

(A) APPLICATION NUMBER: PO9001

(B) FILING DATE: 8 September 1997

(C) APPLICATION NUMBER: PP0398

(D) FILING DATE: 13 November 1997

(viii) ATTORNEY/AGENT INFORMATION:

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(2) INFORMATION FOR SEQ ID NO: 1:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 15 amino acids
- (B) TYPE: amino acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear or cyclic

(ii) MOLECULE TYPE: peptide

(xi) SEQUENCE DESCRIPTION: hGH 177-191

Leu	Arg	Ile	Val	Gln	Cys	Arg	Ser	Val	Glu	Gly	Ser	Cys	Gly	Phe
1				5					10					15

(2) INFORMATION FOR SEQ ID NO: 2:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 17 amino acids
- (B) TYPE: amino acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear or cyclic

(ii) MOLECULE TYPE: peptide

(xi) SEQUENCE DESCRIPTION:

<u>X</u> 1m	Leu	Arg	Ile	Val	Gln	Cys	Arg	Ser	Val	Glu	Gly	Ser	Cys	Gly	Phe	<u>X</u> 2n
1				5					10					15		

(2) INFORMATION FOR SEQ ID NO: 3:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 15 amino acids
- (B) TYPE: amino acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear or cyclic

(ii) MOLECULE TYPE: peptide

(xi) SEQUENCE DESCRIPTION:

<u>Y</u> 1	Leu	Arg	Ile	Val	Gln	Cys	Arg	Ser	Val	Glu	Gly	Ser	Cys	Gly	Phe
1				5					10					15	

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 15 amino acids
- (B) TYPE: amino acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear or cyclic

(xi) SEQUENCE DESCRIPTION:

Leu Arg Ile Val Gln Cys Arg Ser Val Glu Gly Ser Cys Gly Phe Y2
1 5 10 15

(i) SEQUENCE CHARACTERISTICS:
 (A) LENGTH: 15 amino acids
 (B) TYPE: amino acid
 (C) STRANDEDNESS: single
 (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: hGH 177-191 with Cys(Acm) at 182 and 189

Leu Arg Ile Val Gln Cys (Acm) Arg Ser Val Glu Gly Ser Cys (Acm) Gly
Phe
1 5 10 15

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 15 amino acids
- (B) TYPE: amino acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: cyclic

(xi) SEQUENCE DESCRIPTION: hGH 177-191 with Lys at 179

Leu Arg Lys Val Gln Cys Arg Ser Val Glu Gly Ser Cys Gly Phe
1 5 10 15

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(2) INFORMATION FOR SEQ ID NO: 7:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 15 amino acids
- (B) TYPE: amino acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: cyclic

(ii) MOLECULE TYPE: peptide

(xi) SEQUENCE DESCRIPTION: hGH 177-191 with CONH₂ replacing COOH

Leu Arg Ile Val Gln Cys Arg Ser Val Glu Gly Ser Cys Gly Phe -CONH₂
 1 5 10 15

(2) INFORMATION FOR SEQ ID NO: 8:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 15 amino acids
- (B) TYPE: amino acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: cyclic

(ii) MOLECULE TYPE: peptide

(xi) SEQUENCE DESCRIPTION: hGH 177-191 with CH₃CO-HN replacing NH₂

CH₃CO-HN Leu Arg Ile Val Gln Cys Arg Ser Val Glu Gly Ser Cys Gly Phe
 1 5 10 15

(2) INFORMATION FOR SEQ ID NO: 9:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 15 amino acids
- (B) TYPE: amino acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: cyclic

(ii) MOLECULE TYPE: peptide

(xi) SEQUENCE DESCRIPTION: hGH 177-191 with Ala at 183

Leu Arg Ile Val Gln Cys Ala Ser Val Glu Gly Ser Cys Gly Phe
 1 5 10 15

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(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 15 amino acids
- (B) TYPE: amino acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: cyclic

(xi) SEQUENCE DESCRIPTION: hGH 177-191 with Lys at 183

Leu Arg Ile Val Gln Cys Lys Ser Val Glu Gly Ser Cys Gly Phe
1 5 10 15

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 15 amino acids
- (B) TYPE: amino acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: bicyclic

(xi) SEQUENCE DESCRIPTION: hGH 177-191 with Lys at 183 and amide bond between 183 and 186

Leu Arg Ile Val Gln Cys Lys Ser Val Glu Gly Ser Cys Gly Phe
1 5 10 15

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 15 amino acids
- (B) TYPE: amino acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: cyclic

(xi) SEQUENCE DESCRIPTION:hGH 177-191 with H replacing NH₂

H Leu Arg Ile Val Gln Cys Arg Ser Val Glu Gly Ser Cys Gly Phe
1 5 10 15

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 15 amino acids
- (B) TYPE: amino acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: hGH 177-191 with Cys(SH) at 182 and 189

Leu Arg Ile Val Gln Cys(SH) Arg Ser Val Glu Gly Ser Cys(SH) Gly Phe
1 5 10 15

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 15 amino acids
- (B) TYPE: amino acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: cyclic

(xi) SEQUENCE DESCRIPTION: hGH 177-191 with D-Ala at 187 and 190

Leu Arg Ile Val Gln Cys Arg Ser Val Glu D-Ala Ser Cys D-Ala Phe
1 5 10 15

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 15 amino acids
- (B) TYPE: amino acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: cyclic

(xi) SEQUENCE DESCRIPTION: hGH 177-191 with Pen at 182 and 189

Leu Arg Ile Val Gln Pen Arg Ser Val Glu Gly Ser Pen Gly Phe
1 5 10 15

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(2) INFORMATION FOR SEQ ID NO: 16:

- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 15 amino acids
(B) TYPE: amino acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: cyclic

(ii) MOLECULE TYPE: peptide

(xi) SEQUENCE DESCRIPTION: hGH 177-191 with Ala at 191

Leu	Arg	Ile	Val	Gln	Cys	Arg	Ser	Val	Glu	Gly	Ser	Cys	Gly	<u>Ala</u>
1				5					10					15

(2) INFORMATION FOR SEQ ID NO: 17:

- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 15 amino acids
(B) TYPE: amino acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: cyclic

(ii) MOLECULE TYPE: peptide

(xi) SEQUENCE DESCRIPTION: hGH 177-191 with Ala at 190

Leu	Arg	Ile	Val	Gln	Cys	Arg	Ser	Val	Glu	Gly	Ser	Cys	<u>Ala</u>	Phe
1				5					10					15

(2) INFORMATION FOR SEQ ID NO: 18:

- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 15 amino acids
(B) TYPE: amino acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: cyclic

(ii) MOLECULE TYPE: peptide

(xi) SEQUENCE DESCRIPTION: hGH 177-191 with Ala at 178

Leu	<u>Ala</u>	Ile	Val	Gln	Cys	Arg	Ser	Val	Glu	Gly	Ser	Cys	Gly	Phe
1				5					10					15

0000240445080560

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 16 amino acids
- (B) TYPE: amino acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: cyclic

(ii) MOLECULE TYPE: peptide

(xi) SEQUENCE DESCRIPTION: Tyr - hGH 177-191

Tyr Leu Arg Ile Val Gln Cys Arg Ser Val Glu Gly Ser Cys Gly Phe
1 5 10 15

(2) INFORMATION FOR SEQ ID NO: 20:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 16 amino acids
- (B) TYPE: amino acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: cyclic

(ii) MOLECULE TYPE: peptide

(xi) SEQUENCE DESCRIPTION: Lys - hGH 177-191

Lys Leu Arg Ile Val Gln Cys Arg Ser Val Glu Gly Ser Cys Gly Phe
1 5 10 15

(2) INFORMATION FOR SEQ ID NO: 21:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 15 amino acids
- (B) TYPE: amino acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: cyclic

(ii) MOLECULE TYPE: peptide

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(xi) SEQUENCE DESCRIPTION: hGH 177-191 with Lys at 178

Leu Lys Ile Val Gln Cys Arg Ser Val Glu Gly Ser Cys Gly Phe
1 5 10 15

(2) INFORMATION FOR SEQ ID NO: 22:

- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 15 amino acids
(B) TYPE: amino acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: cyclic

(ii) MOLECULE TYPE: peptide

(xi) SEQUENCE DESCRIPTION: hGH 177-191 with Ala at 177

Ala Arg Ile Val Gln Cys Arg Ser Val Glu Gly Ser Cys Gly Phe
1 5 10 15

(2) INFORMATION FOR SEQ ID NO: 23:

- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 15 amino acids
(B) TYPE: amino acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: cyclic

(ii) MOLECULE TYPE: peptide

(xi) SEQUENCE DESCRIPTION: hGH 177-191 with Ala at 179

Leu Arg Ala Val Gln Cys Arg Ser Val Glu Gly Ser Cys Gly Phe
1 5 10 15

(2) INFORMATION FOR SEQ ID NO: 24:

- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 15 amino acids
(B) TYPE: amino acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: cyclic

(ii) MOLECULE TYPE: peptide

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(xi) SEQUENCE DESCRIPTION: hGH 177-191 with Ala at 180

Leu Arg Ile Ala Gln Cys Arg Ser Val Glu Gly Ser Cys Gly Phe
1 5 10 15

(2) INFORMATION FOR SEQ ID NO: 25:

- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 15 amino acids
(B) TYPE: amino acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: cyclic

(ii) MOLECULE TYPE: peptide

(xi) SEQUENCE DESCRIPTION: hGH 177-191 with Ala at 181

Leu Arg Ile Val Ala Cys Arg Ser Val Glu Gly Ser Cys Gly Phe
1 5 10 15

(2) INFORMATION FOR SEQ ID NO: 26:

- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 15 amino acids
(B) TYPE: amino acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(xi) SEQUENCE DESCRIPTION: hGH 177-191 with Ala at 182

Leu Arg Ile Val Gln Ala Arg Ser Val Glu Gly Ser Cys Gly Phe
1 5 10 15

(2) INFORMATION FOR SEQ ID NO: 27:

- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 15 amino acids
(B) TYPE: amino acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: cyclic

(ii) MOLECULE TYPE: peptide

Leu Arg Ile Val Gln Cys Arg Ala Val Glu Gly Ser Cys Gly Phe
1 5 10 15

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 15 amino acids
- (B) TYPE: amino acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: cyclic

(xi) SEQUENCE DESCRIPTION: hGH 177-191 with Ala at 185

Leu Arg Ile Val Gln Cys Arg Ser Ala Glu Gly Ser Cys Gly Phe
1 5 10 15

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 15 amino acids
- (B) TYPE: amino acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: cyclic

(xi) SEQUENCE DESCRIPTION: hGH 177-191 with Ala at 186

Leu Arg Ile Val Gln Cys Arg Ser Val Ala Gly Ser Cys Gly Phe
1 5 10 15

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 15 amino acids
- (B) TYPE: amino acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: cyclic

(ii) MOLECULE TYPE: peptide

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(xi) SEQUENCE DESCRIPTION: hGH 177-191 with Ala at 187

Leu Arg Ile Val Gln Cys Arg Ser Val Glu Ala Ser Cys Gly Phe
1 5 10 15

(2) INFORMATION FOR SEQ ID NO: 31:

- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 15 amino acids
(B) TYPE: amino acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: cyclic

(ii) MOLECULE TYPE: peptide

(xi) SEQUENCE DESCRIPTION: hGH 177-191 with Ala at 188

Leu Arg Ile Val Gln Cys Arg Ser Val Glu Gly Ala Cys Gly Phe
1 5 10 15

(2) INFORMATION FOR SEQ ID NO: 32:

- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 15 amino acids
(B) TYPE: amino acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

(xi) SEQUENCE DESCRIPTION: hGH 177-191 with Ala at 189

Leu Arg Ile Val Gln Cys Arg Ser Val Glu Gly Ser Ala Gly Phe
1 5 10 15

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(2) INFORMATION FOR SEQ ID NO: 33:

- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 17 amino acids
(B) TYPE: amino acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: cyclic

(ii) MOLECULE TYPE: peptide

(xi) SEQUENCE DESCRIPTION: Lys-Lys- hGH 177-191

Lys Lys Leu Arg Ile Val Gln Cys Arg Ser Val Glu Gly Ser Cys Gly Phe
1 5 10 15